

26352U sequence listing.ST25.txt
SEQUENCE LISTING

<110> Japan Science and Technology Agency

<120> Gene participating in the synthesis of brassinosteroid

<130> 26352U (PS03-311PCT)

<160> 6

<170> PatentIn version 3.1

<210> 1

<211> 1473

<212> DNA

<213> Arabidopsis thaliana

<400> 1

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atcggtgaaa ccatcgagtt cgtctcttct gcttactcag accgtcctga gagtttcatg      240
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<210> 2
 <211> 490
 <212> PRT
 <213> Arabidopsis thaliana

<400> 2

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20         25         30

Ser Lys Lys Lys Leu Asn Asp His His Val Thr Ser Gln Ser His Gly
35         40         45

Pro Lys Phe Pro His Gly Ser Leu Gly Trp Pro Val Ile Gly Glu Thr
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Ile Glu Phe Val Ser Ser Ala Tyr Ser Asp Arg Pro Glu Ser Phe Met
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Asp Lys Arg Arg Leu Met Tyr Gly Arg Val Phe Lys Ser His Ile Phe
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Gly Thr Ala Thr Ile Val Ser Thr Asp Ala Glu Val Asn Arg Ala Val
100        105        110

Leu Gln Ser Asp Ser Thr Ala Phe Val Pro Phe Tyr Pro Lys Thr Val
115        120        125

Arg Glu Leu Met Gly Lys Ser Ser Ile Leu Leu Ile Asn Gly Ser Leu
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His Arg Arg Phe His Gly Leu Val Gly Ser Phe Leu Lys Ser Pro Leu
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Leu Lys Ala Gln Ile Val Arg Asp Met His Lys Phe Leu Ser Glu Ser
165        170        175

Met Asp Leu Trp Ser Glu Asp Gln Pro Val Leu Leu Gln Asp Val Ser
180        185        190

Lys Thr Val Ala Phe Lys Val Leu Ala Lys Ala Leu Ile Ser Val Glu
195        200        205

Lys Gly Glu Asp Leu Glu Glu Leu Lys Arg Glu Phe Glu Asn Phe Ile
210        215        220

Ser Gly Leu Met Ser Leu Pro Ile Asn Phe Pro Gly Thr Gln Leu His
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Ile Glu Gly Lys Ile Arg Lys Thr Lys Asn Lys Glu Glu Asp Asp Val
260 265 270

Ile Ala Lys Asp Val Val Asp Val Leu Leu Lys Asp Ser Ser Glu His
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Leu Thr His Asn Leu Ile Ala Asn Asn Met Ile Asp Met Met Ile Pro
290 295 300

Gly His Asp Ser Val Pro Val Leu Ile Thr Leu Ala Val Lys Phe Leu
305 310 315 320

Ser Asp Ser Pro Ala Ala Leu Asn Leu Leu Thr Lys Asn Met Lys Leu
325 330 335

Lys Ser Leu Lys Glu Leu Thr Gly Glu Pro Leu Tyr Trp Asn Asp Tyr
340 345 350

Leu Ser Leu Pro Leu Thr Gln Lys Val Ile Thr Glu Thr Leu Arg Met
355 360 365

Gly Asn Val Ile Ile Gly Val Met Arg Lys Ala Met Lys Asp Val Glu
370 375 380

Ile Lys Gly Tyr Val Ile Pro Lys Gly Trp Cys Phe Leu Ala Tyr Leu
385 390 395 400

Arg Ser Val His Leu Asp Glu Ala Tyr Tyr Glu Ser Pro Tyr Lys Phe
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Asn Pro Trp Arg Trp Gln Glu Arg Asp Met Asn Thr Ser Ser Phe Ser
420 425 430

Pro Phe Gly Gly Gly Gln Arg Leu Cys Pro Gly Leu Asp Leu Ala Arg
435 440 445

Leu Glu Thr Ser Val Phe Leu His His Leu Val Thr Arg Phe Arg Trp
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Asn Lys Leu Pro Ile Trp Ile Lys Arg Ile
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<210> 3
<211> 1934
<212> DNA
<213> Arabidopsis thaliana

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<220>
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 <222> (1748)..(1748)
 <223> n means A, C, G or T.

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tttgtttgtc atgtcaaatt ataagcgttg gttaggttgc ccctttctct tttatttatc 1860
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atatagcggc cggg 1934

<210> 4
<211> 524
<212> PRT
<213> Arabidopsis thaliana
<400> 4

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20 25 30

Leu Thr Ala Gly Ile Leu Leu Arg Pro Trp Leu Trp Phe Arg Leu Arg
35 40 45

Asn Ser Lys Thr Lys Asp Gly Asp Glu Glu Glu Asp Asn Glu Glu Lys
50 55 60

Lys Lys Gly Met Ile Pro Asn Gly Ser Leu Gly Trp Pro Val Ile Gly
65 70 75 80

Glu Thr Leu Asn Phe Ile Ala Cys Gly Tyr Ser Ser Arg Pro Val Thr
85 90 95

Phe Met Asp Lys Arg Lys Ser Leu Tyr Gly Lys Val Phe Lys Thr Asn
100 105 110

Ile Ile Gly Thr Pro Ile Ile Ile Ser Thr Asp Ala Glu Val Asn Lys
115 120 125

Val Val Leu Gln Asn His Gly Asn Thr Phe Val Pro Ala Tyr Pro Lys
130 135 140

Ser Ile Thr Glu Leu Leu Gly Glu Asn Ser Ile Leu Ser Ile Asn Gly
145 150 155 160

Pro His Gln Lys Arg Leu His Thr Leu Ile Gly Ala Phe Leu Arg Ser
165 170 175

Pro His Leu Lys Asp Arg Ile Thr Arg Asp Ile Glu Ala Ser Val Val
180 185 190

Leu Thr Leu Ala Ser Trp Ala Gln Leu Pro Leu Val His Val Gln Asp
195 200 205

Glu Ile Lys Lys Met Thr Phe Glu Ile Leu Val Lys Val Leu Met Ser
210 215 220

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Thr Ser Pro Gly Glu Asp Met Asn Ile Leu Lys Leu Glu Phe Glu Glu
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Phe Ile Lys Gly Leu Ile Cys Ile Pro Ile Lys Phe Pro Gly Thr Arg
245 250 255

Leu Tyr Lys Ser Leu Lys Ala Lys Glu Arg Leu Ile Lys Met Val Lys
260 265 270

Lys Val Val Glu Glu Arg Gln Val Ala Met Thr Thr Thr Ser Pro Ala
275 280 285

Asn Asp Val Val Asp Val Leu Leu Arg Asp Gly Gly Asp Ser Glu Lys
290 295 300

Gln Ser Gln Pro Ser Asp Phe Val Ser Gly Lys Ile Val Glu Met Met
305 310 315 320

Ile Pro Gly Glu Glu Thr Met Pro Thr Ala Met Thr Leu Ala Val Lys
325 330 335

Phe Leu Ser Asp Asn Pro Val Ala Leu Ala Lys Leu Val Glu Glu Asn
340 345 350

Met Glu Met Lys Arg Arg Lys Leu Glu Leu Gly Glu Glu Tyr Lys Trp
355 360 365

Thr Asp Tyr Met Ser Leu Ser Phe Thr Gln Asn Val Ile Asn Glu Thr
370 375 380

Leu Arg Met Ala Asn Ile Ile Asn Gly Val Trp Arg Lys Ala Leu Lys
385 390 395 400

Asp Val Glu Ile Lys Gly Tyr Leu Ile Pro Lys Gly Trp Cys Val Leu
405 410 415

Ala Ser Phe Ile Ser Val His Met Asp Glu Asp Ile Tyr Asp Asn Pro
420 425 430

Tyr Gln Phe Asp Pro Trp Arg Trp Asp Arg Ile Asn Gly Ser Ala Asn
435 440 445

Ser Ser Ile Cys Phe Thr Pro Phe Gly Gly Gly Gln Arg Leu Cys Pro
450 455 460

Gly Leu Glu Leu Ser Lys Leu Glu Ile Ser Ile Phe Leu His His Leu
465 470 475 480

Val Thr Arg Tyr Ser Trp Thr Ala Glu Glu Asp Glu Ile Val Ser Phe
485 490 495

26352U sequence listing.ST25.txt

Pro Thr Val Lys Met Lys Arg Arg Leu Pro Ile Arg Val Ala Thr Val
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Asp Asp Ser Ala Ser Pro Ile Ser Leu Glu Asp His
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<210> 5
 <211> 20
 <212> DNA
 <213> Artificial sequence

<220>
 <223> PCR amplification primer

<400> 5
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<210> 6
 <211> 21
 <212> DNA
 <213> Artificial sequence

<220>
 <223> PCR amplification primer

<400> 6
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